

1/ /

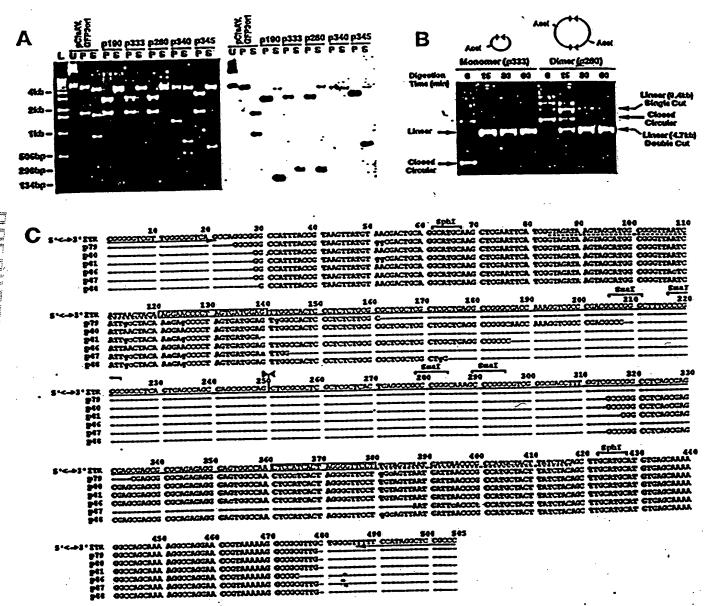


Figure &

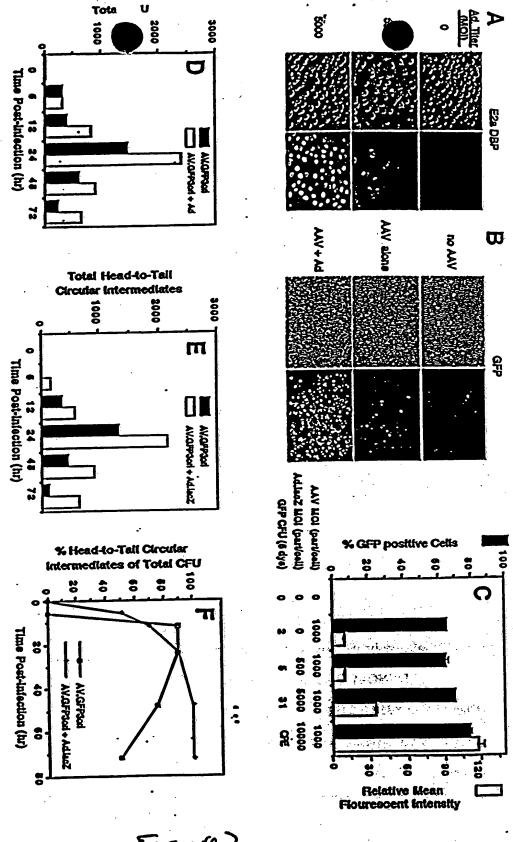


Figure 3

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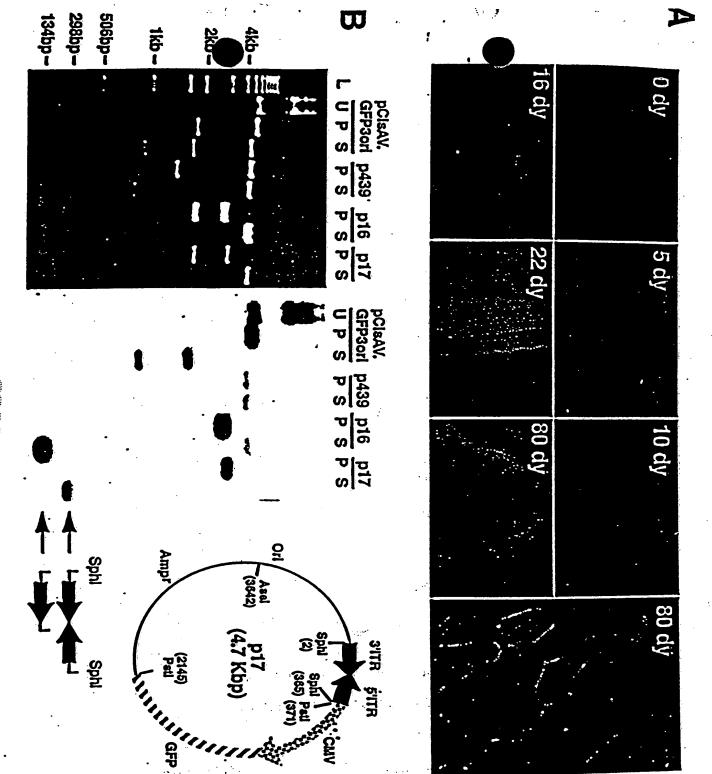


Figure 4

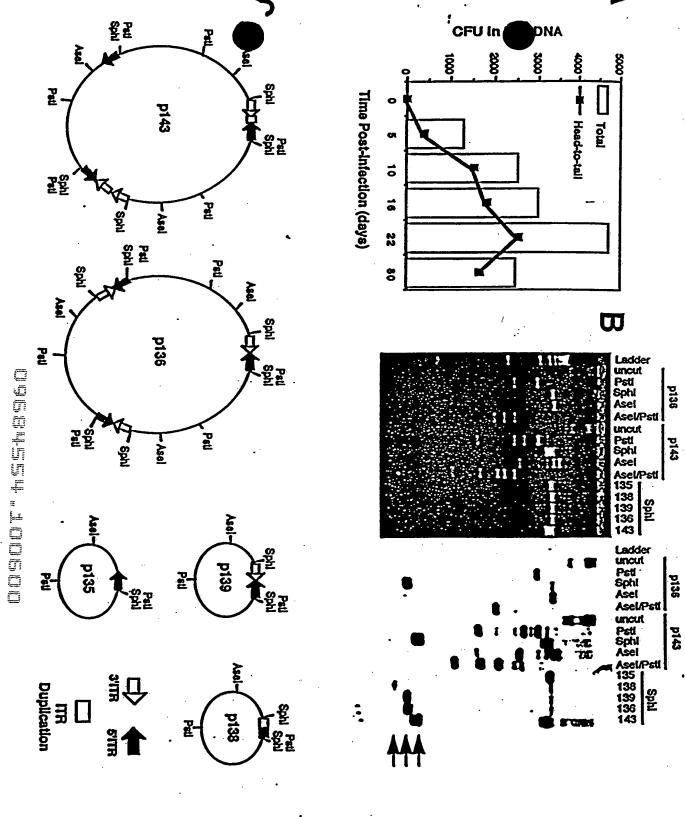
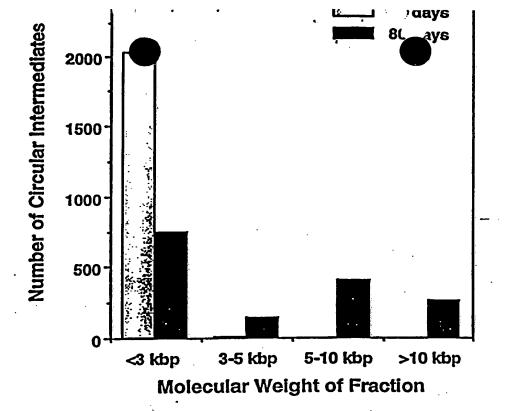
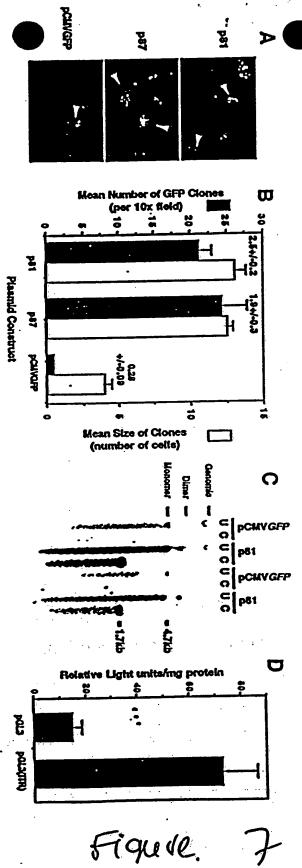


Figure 5

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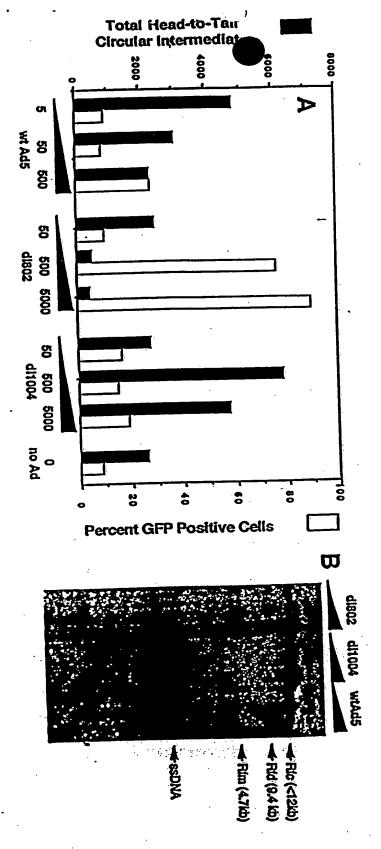


Figure 8

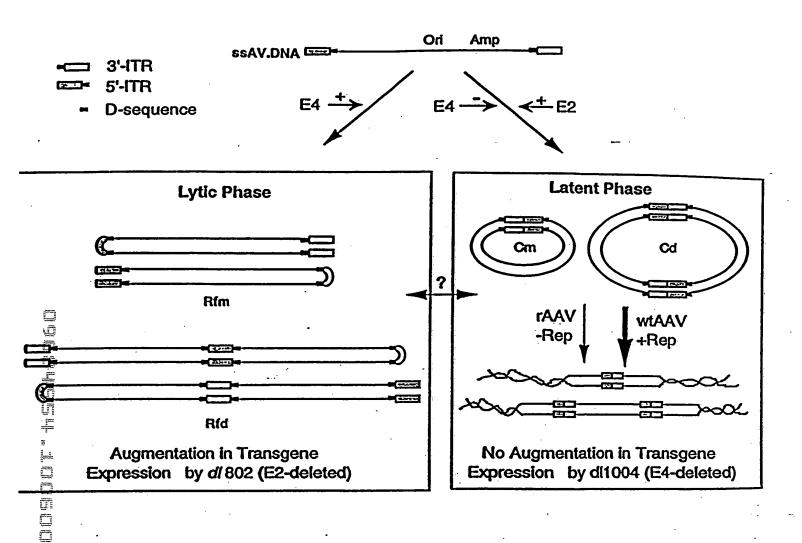


Figure 9

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- 4	-

43.					
10	20	30	40	50	60
GCATGCAAGC	TGTAGATAAG	TAGCATGGCG	GGTTAATCAT	TAACTACAAG	GAACCCCTAG
CGTACGTTCG	ACATCTATTC	ATCGTACCGC	CCAATTAGTA	ATTGATGTTC	CTTGGGGATC
·		•	•		
. 70	80	90	100	110	120
TGATGGAGTT	GGCCACTCCC	TCTCTGCGCG	CTCGCTCGCT	CACTGAGGCC	
	CCGGTGAGGG				
130	140	150	160	170	180
AGGTCGCCCG	ACGCCCGGGC	TTTGCCCGGG	CGGCCTCAGT	GAGCGAGCGA	GCGCGCAGAG
TCCAGCGGGC	TGCGGGCCCG	AAACGGGCCC	GCCGGAGTCA	CTCGCTCGCT	CGCGCGTCTC
	•			,	
190	200	210	220	230	240
AGGGAGTGGC	CAACTCCATC	ACTAGGGGTT	CCTTGTAGTT	AATGATTAAC.	CCGCCATGCT
TCCCTCACCG	GTTGAGGTAG	TGATCCCCAA	GGAACATCAA	TTACTAATTG	GGCGGTACGA
:	•	,			
250	260	270	280		
ACTTATCTAC	CGATGAATTC	GAGCTTGCAT	<u>GC</u>		,
TGAATAGATG	GCTACTTAAG	CTCGAACGTA	C G		
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ui In 10	20	30	40	50	60
*** · · ·	TGTAGATAAG				
CCTACCTTCG			•		
	illicite in the second	111001110000	CCIMITAGIA	ALIGAIGITC	CIIGGGAIC
70	80	90	100	110	120
TGATGGAGTT				110	120
ACTACCTCAA					
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	•				
130	140	150	160	170	180
TOGCTOGCTC		150 GGCGACCAAA	160		180
TOGOTOGOTO	ACTGAGGCCG	GGCGACCAAA	GGTCGCCCGA	GCCCGGCTT	TGCCCGGGCG
TOGOTOGOTO		GGCGACCAAA	GGTCGCCCGA	GCCCGGCTT	TGCCCGGGCG
TOGOTOGOTO	ACTGAGGCCG	GGCGACCAAA	GGTCGCCCGA	GCCCGGGCTT CGGGCCCGAA	TGCCCGGGCG ACGGGCCCGC
TOGOTOGOTO AGCGAGOGAG	ACTGAGGCCG TGACTCCGGC	GGCGACCAAA CCGCTGGTTT 210	GGTCGCCCGA CCAGCGGGCT 220	GCCCGGCTT CGGGCCCGAA 230	TGCCCGGGGG ACGGGCCCGC
TOGOTOGOTO AGOGAGOGAG 190 GCOTOAGTGA	ACTGAGGCCG TGACTCCGGC 200 GCGAGCGCGC	GGCGACCAAA CCGCTGGTTT 210 GCGCAGAGAG	GGTCGCCCGA CCAGCGGGCT 220 GGAGTGGCCA	GCCCGGCTT CGGGCCCGAA 230 ACTCCATCAC	TGCCCGGCG ACGGGCCCGC 240 TAGGGGTTCC
TOGOTOGOTO AGOGAGOGAG 190 GCOTOAGTGA	ACTGAGGCCG TGACTCCGGC 200	GGCGACCAAA CCGCTGGTTT 210 GCGCAGAGAG	GGTCGCCCGA CCAGCGGGCT 220 GGAGTGGCCA	GCCCGGCTT CGGGCCCGAA 230 ACTCCATCAC	TGCCCGGCG ACGGGCCCGC 240 TAGGGGTTCC
TOGOTOGOTO AGOGAGOGAG 190 GCOTOAGTGA	ACTGAGGCCG TGACTCCGGC 200 GCGAGCGCGC	GGCGACCAAA CCGCTGGTTT 210 GCGCAGAGAG	GGTCGCCCGA CCAGCGGGCT 220 GGAGTGGCCA CCTCACCGGT	GCCCGGCTT CGGGCCCGAA 230 ACTCCATCAC	TGCCCGGGCG ACGGGCCCGC 240 TAGGGGTTCC ATCCCCAAGG
TOGCTCGCTC AGCGAGCGAG 190 GCCTCAGTGA CGGAGTCACT	ACTGAGGCCG TGACTCCGGC 200 GCGAGCGCGC CGCTCGCGCG	GGCGACCAAA CCGCTGGTTT 210 GCGCAGAGAG CGCGTCTCTC	GGTCGCCCGA CCAGCGGGCT 220 GGAGTGGCCA CCTCACCGGT	GCCCGGGCTT CGGGCCCGAA 230 ACTCCATCAC TGAGGTAGTG	TGCCCGGGCG ACGGGCCCGC 240 TAGGGGTTCC ATCCCCAAGG

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10	20	30	40	50	60
GCATGCAAGC	TGTAGATAAG	TAGCATGGCG	GGTTAATCAT	TAACTACAAG	GAACCCCTAG
CGTACGTTCG	ACATCTATTC	ATCGTACCGC	CCAATTAGTA	ATTGATGTTC	CTTGGGGATC
70	80	90	-		120
TGATGGAGTT	GGCCACTCCC	TCTCTGCGCG	CICGCICGCI	CACTGAGGCC	GGGCGACCAA
ACTACCTCAA	CCGGTGAGGG	AGAGACGCGC	GAGCGAGCGA	GTGACTCCGG	CCCGCTGGTT
130	. 140	150	160	170	180
				GAGCGAGCGA	
TECAGCGGGC				CTCGCTCGCT	
0 - 190	200	210	220	230	240
AGGGAGTGGC	CAACTCCATC	ACTAGGGGTT	CCTTGTAGTT	AATGATTAAC	CCGCCATGCT
TOCCTCACCG	GTTGAGGTAG	TGATCCCCAA	GGAACATCAA	TTACTAATTG	GGCGGTACGA
250	260	270	280		
ACTTATCTAC	CGATGAATTC	GAGCTTGCAT	GC		-
TGAATAGATG	GCTACTTAAG	CTCGAACGTA	<u>CG</u>		
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		10 .	20.	30	40	50	
			TARMAD AMON	TACCATGGCG	GGTTAATCAT	TAACTACAAG	50
P81			COMPARAGE COMPANY	CONTRACTOR TO A T	CCLIMITOR	TIME TIME	
p 79	1	GCATGCAAGC	TGTLGATAAG	TAGCATGGCG	GGTTAATCAT	TAACTACAAG	50
p1202	1	GCATGCAAGC	Idinguina				
		60	70	80	90	100	•
				GGCCACTCCC	TCTCTGCGCG	CTCGCTCGCT	100
P81							
p79	51	GAACCCCTAG	TGATGGAGTT	GGCCACTCCC	TCTCTGCGCG	CTCGCTCGCT	100
p1202	51						
		110	120	130	140	150	
	- 01		00000			@@@###	150
				Tryprit 12 11.	MCIGUGGCCG	00000	_
p79	101	CACTGAGGCC	GGGCG			ACCAAAG	1:50
[™] p1202	101	CACIGAGGG					
		160	170	180	190	200	
-: Il -01	151	CTCCCCGAC	GCCCGGCTI	TGCCCGGGCG	GCCTCAGTGA	GCGAGCGAGC	200
P81 P79							
* p /y	151	GTCGCCCGA(GCCCGGCTT	TGgtCGccCG	GCCTCAGTGA	GCGAGCGAGC	200
p1202	191	020000			_		
Adhanne Managari Managari Managari Managari Managari Managari		210	220	230	240	,	
P81 p79	201	GCGCAGAGA		A CONTRACTOR A	TAGGGGTTC	TTGTAGTTAA	250
_ p/3	201	GCGCAGAGA	G GGAGTGGCC	A ACTCCATCA	C TAGGGGTTC	TTGTAGTTAA	250
p1202	20.				•	_	
•		26	0 27	0 28	0 29		
P81	25	1 TGATTAACC	C GCCATGCTA	C TTATCTACC	G ATGAATICG	A GCTTGCATGO	300
p79				רי אויאיוייף י	C VIGURITO	Tr Core	_
P13			C GCCATGCTA	C TTATCTACC	G ATGAATTCG	A GCTTGCATG	_ 500

Figure 11

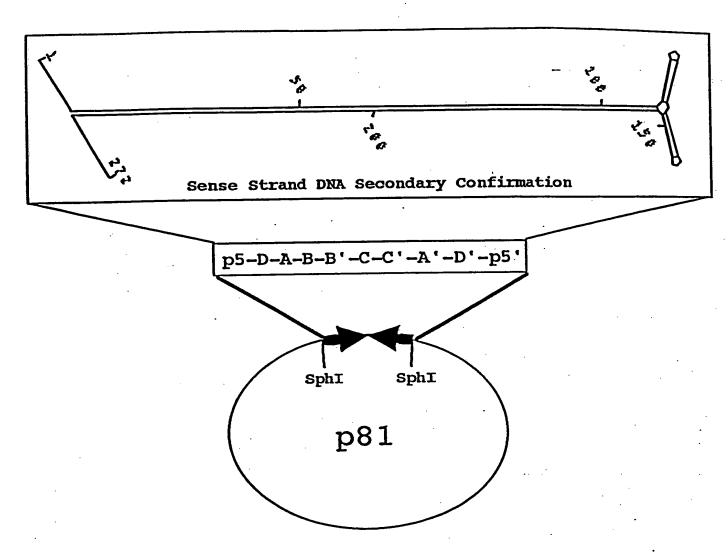


Figure (2A

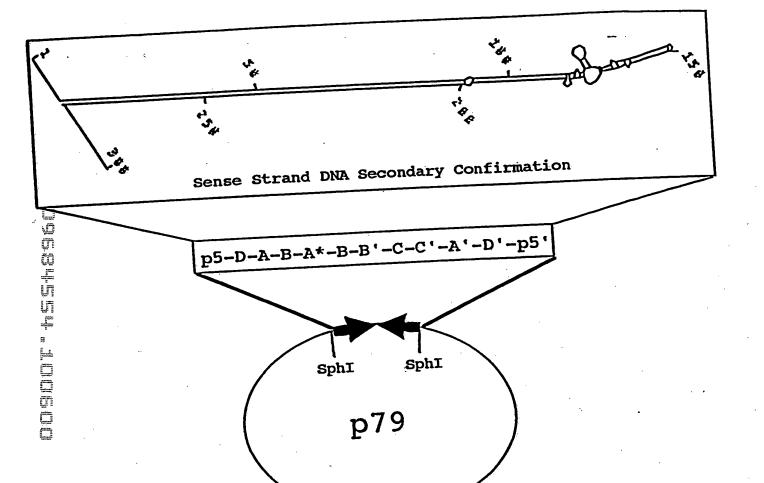
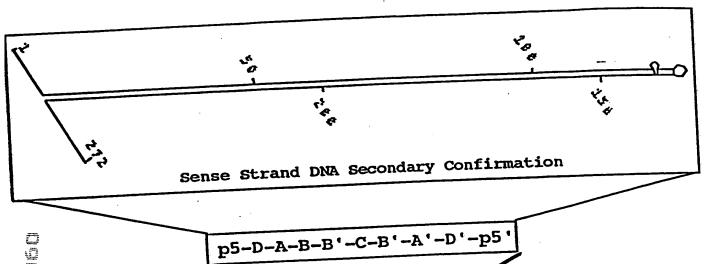
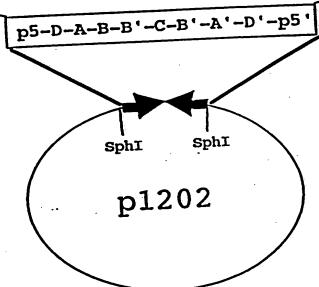


Figure 12B





: Figure 12 C

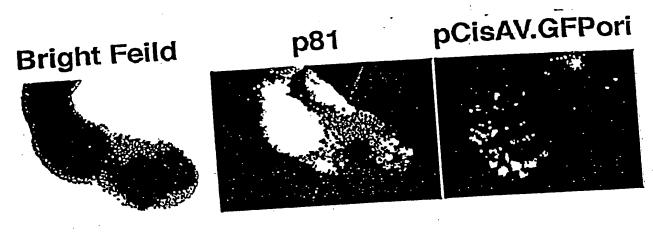


Figure 13

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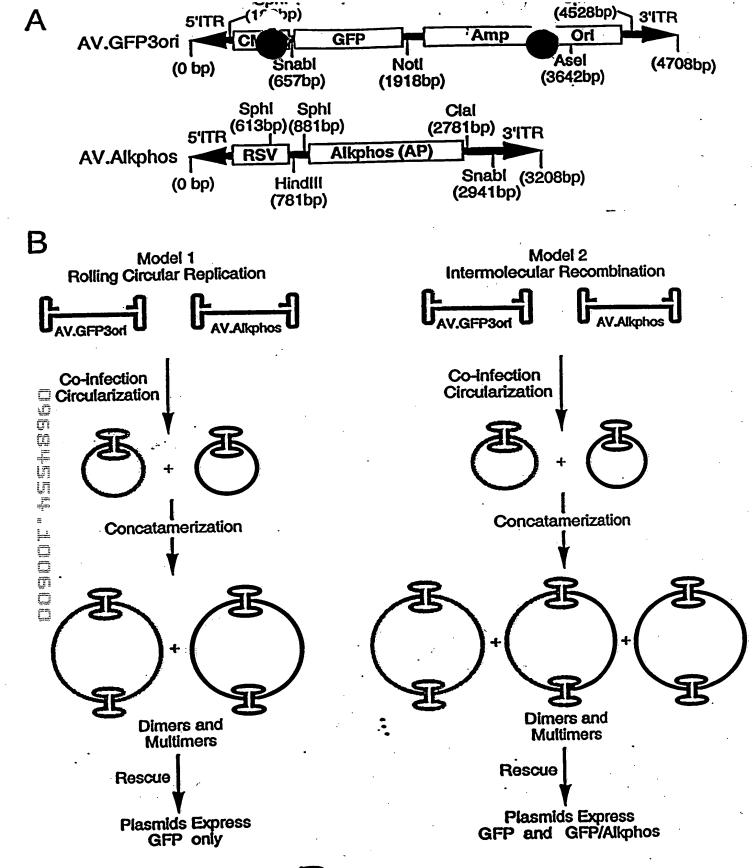


Figure 14

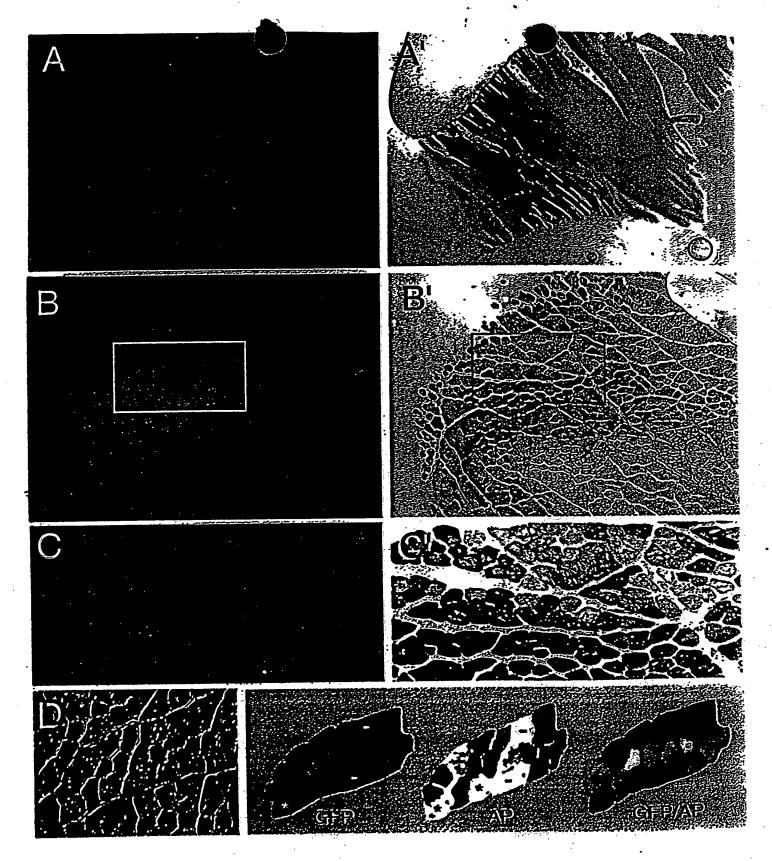


Figure 15

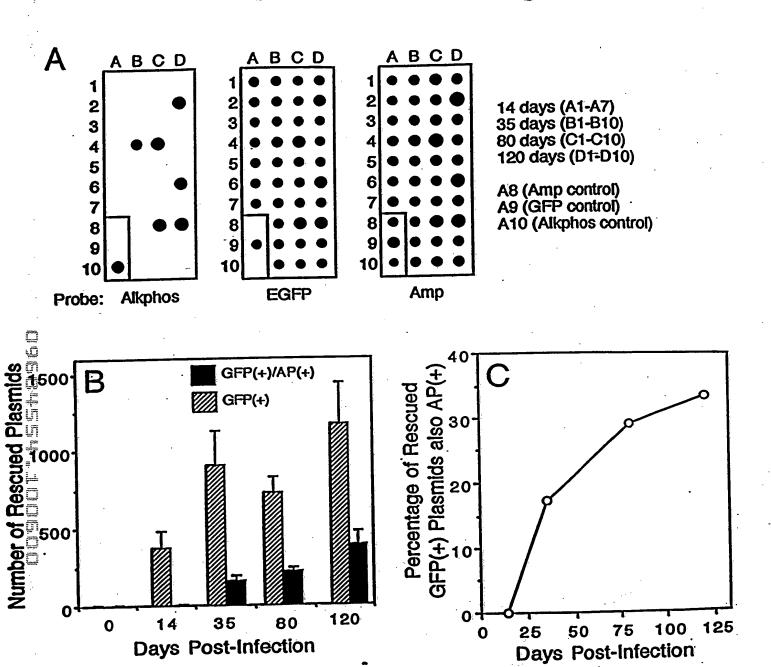
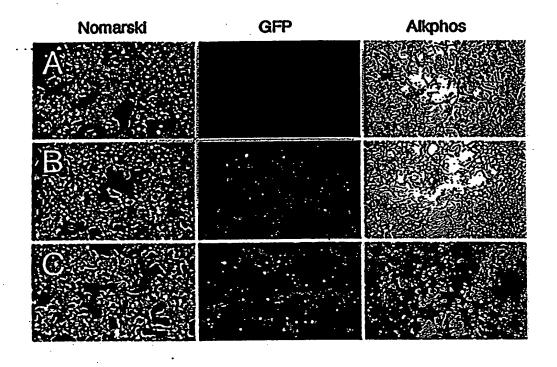
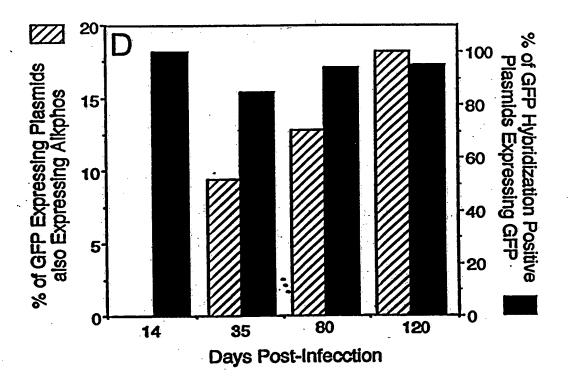


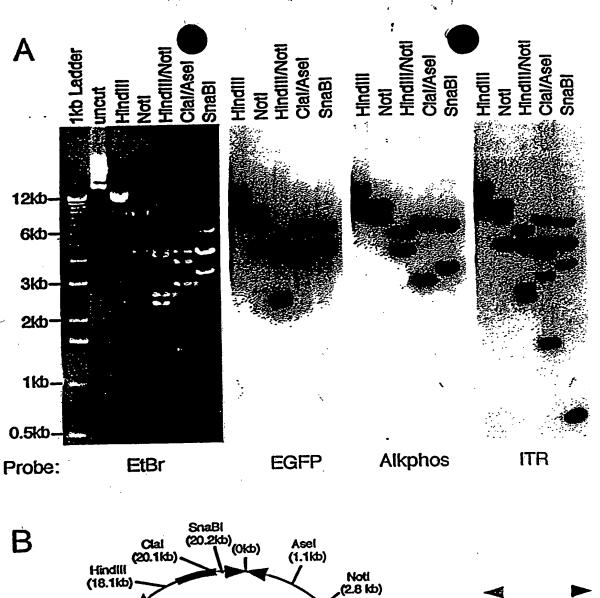
Figure 16

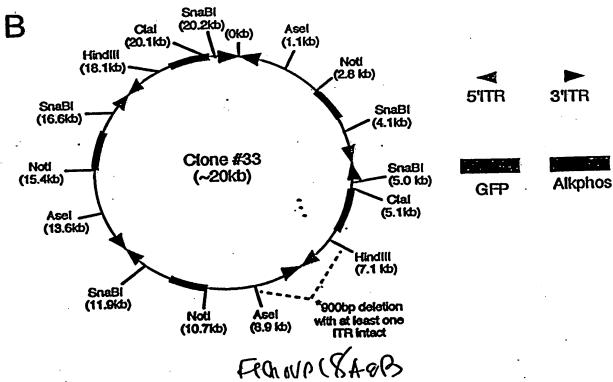
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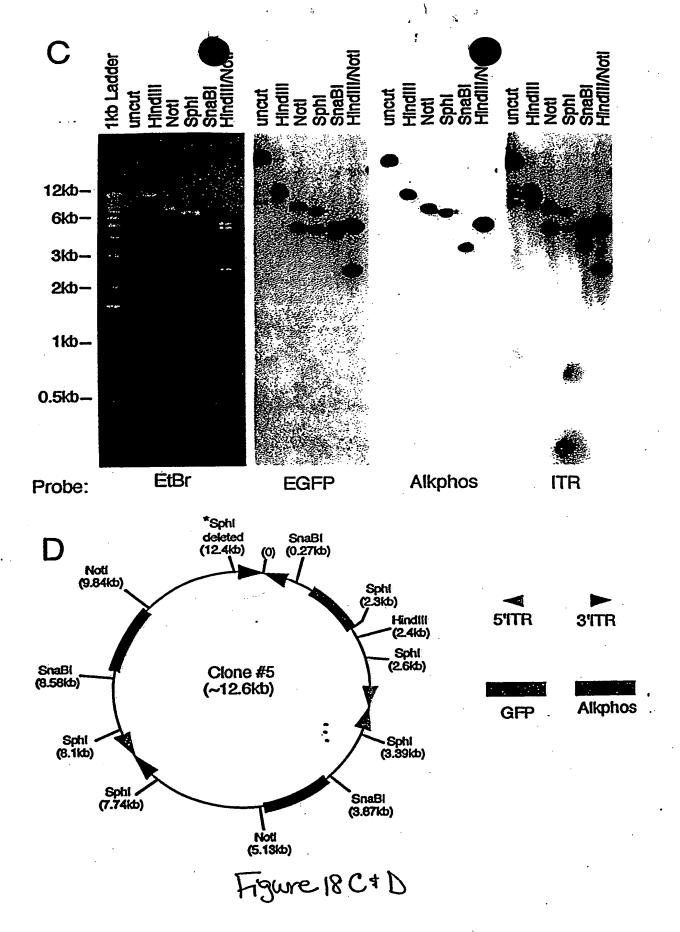


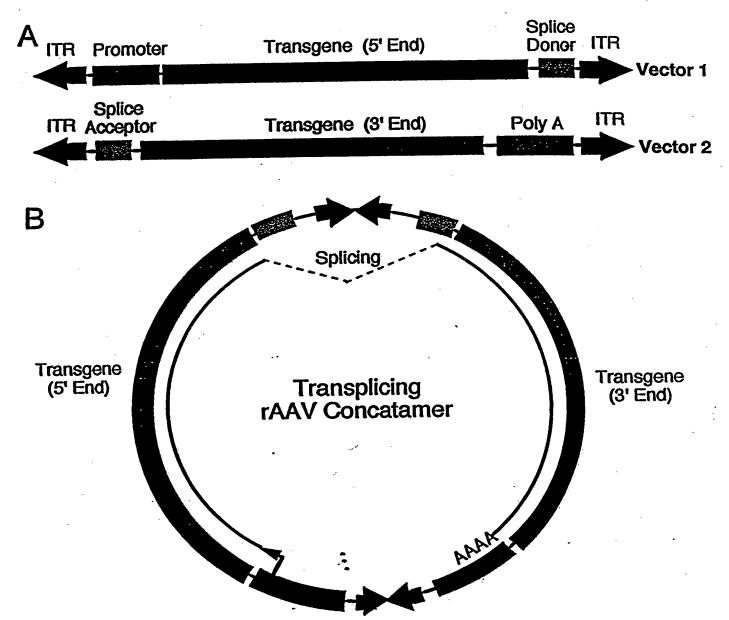


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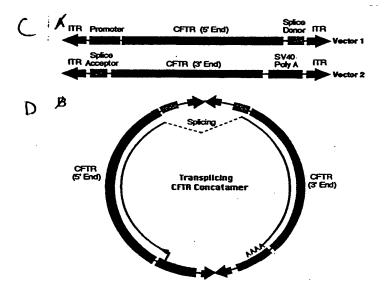
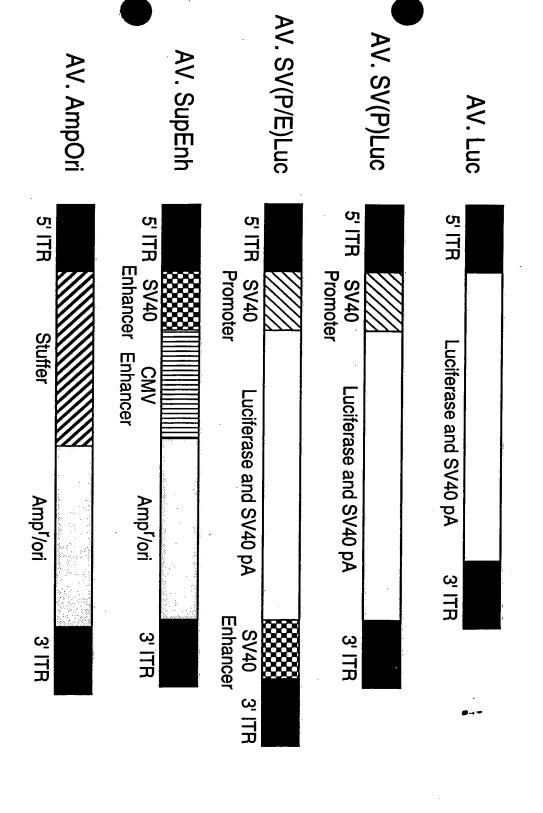
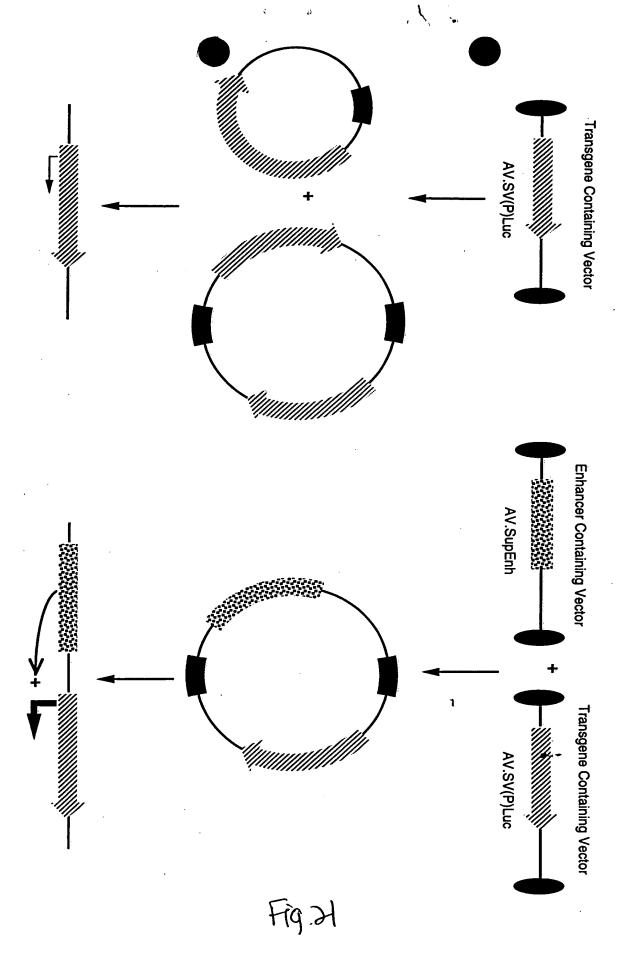


Fig 19 Candb



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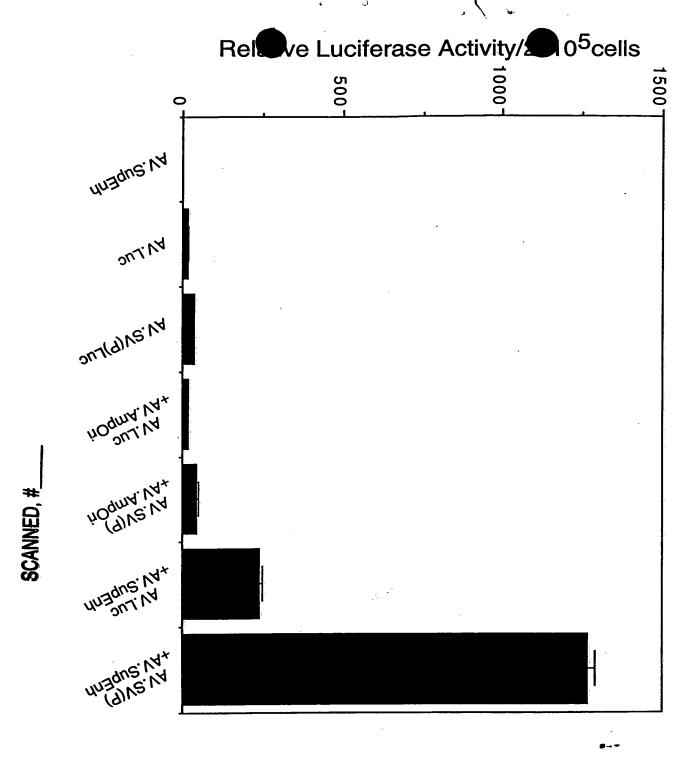
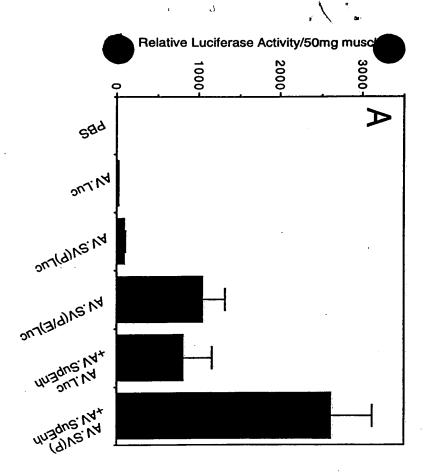


Fig. 22



Relative Luciferase Activity/50mg muscle

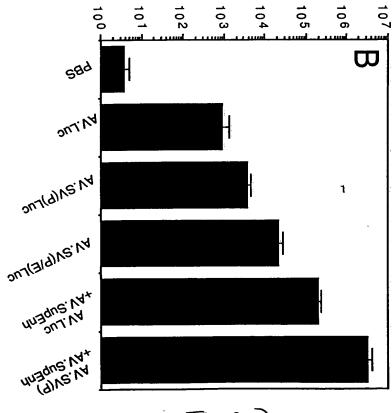
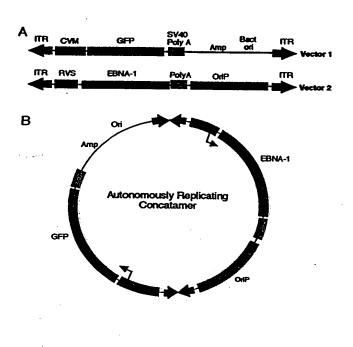


Fig. 23



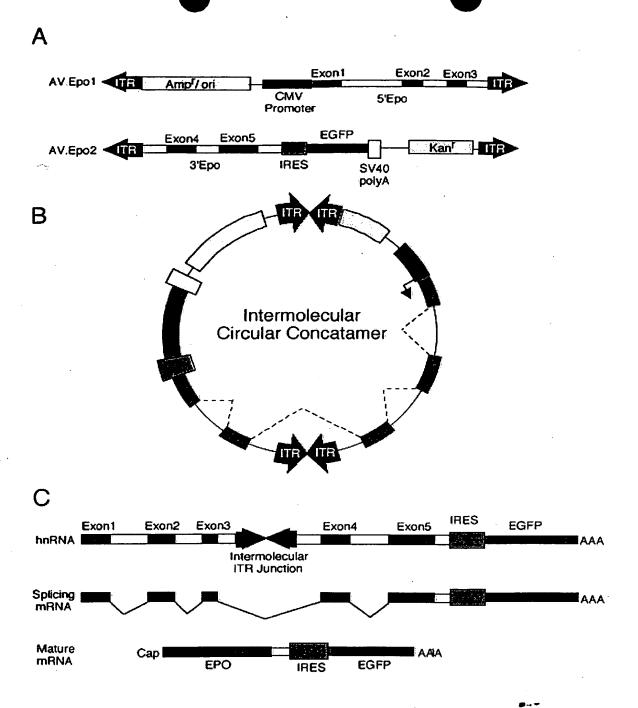


Fig.25

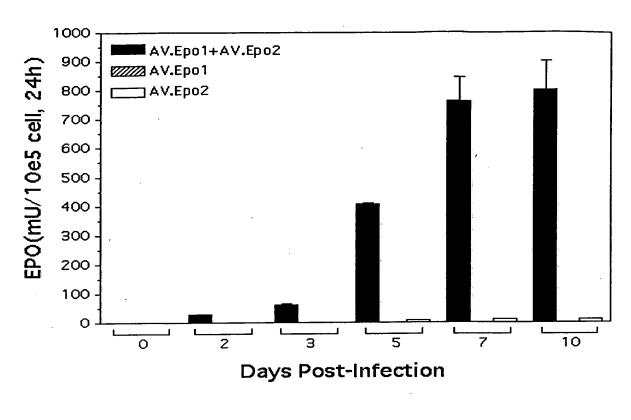


Fig 26

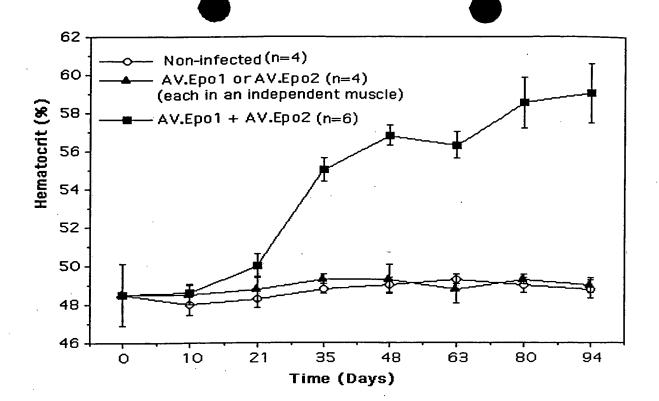


Fig 2?